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ID      091A69;          PRELIMINARY;          PRT;    197 AA.
DB       01-OCT-2000 (TEMBUREL_15; Created)
DB       01-JUN-2001 (TEMBUREL_15; Last sequence update)
DB       01-JUN-2001 (TEMBUREL_17; Last annotation update)
DB       TIRINGO A; Accession 1 (Phantom).
CC       Gallus gallus (Chicken)
CC       Archaeosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
CC       Gallus.
CC       NCBI_TaxID=9031;
RN       11)
RP       SEQUENCE FROM N.A.
RC       STRAIN-BRED FAYOUMT; TTSUSE-SPLEEN;
RA       Zhou H., Lamont S.J.;
RF       specific genes1, on among chicken lines and mammalian species in
RL       submitted (JAN-2000) to the EMBL/Genbank/DBD databases.
DR       EMBL: AF221083; AAF5320.1;
DR       InterPro: IPR001370; BIR.
DR       Pfam: PF00653; BIR. 2.
DR       SMART: SM00283; BIR. 2.
DR       PROSITE: PS01282; BIR_REPEAT_1; 2.
DR       PROSITE: PS50143; BIR_REPEAT_2; 2.
DR       NON_TER
ET       197
SQ       SOURCE     197 AA; 22602 MW; D7923DABCF623E3A CRR64;

Query Match
Best Local Similarity 98.3%; Score 290; DB 13; Length 197;
Matches 46; Conservative 1; Mismatches 1; Indels 0; Gaps
DY       -1 PEGLASAGFYVGRNDYVCFCDCGGILRCWESGDDFWERAKKVPFCE 48
DB       135 FEQLDAAGFYVGRNDYVCFCDCGGILRCWESGDDFWERAKKVPFCE 182
RESULT   3
057319
ID      057319;          PRELIMINARY;          PRT;    610 AA.
AC       057319;
DT       01-JUN-1998 (TEMBUREL_06; Created)
DT       01-JUN-1998 (TEMBUREL_06; Last sequence update)
DT       01-JUN-2001 (TEMBUREL_17; Last annotation update)
DE       D1-100001 (Tembeurel_17, last annotation update)
DE       D1-100001 OF NEOPTOSIS PROTEIN 1 (IAP) (INHIBITOR OF T CELL APOPTOSIS
DE       PROTEIN).
OS       Gallus gallus (Chicken).
GN       Iap1.
OC       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC       Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
OC       Gallus.
OC       NCBI_TaxID=9031;
RN       11)
RP       SEQUENCE FROM N.A.
RC       STRAIN-BRED FAYOUMT; TTSUSE-SPLEEN;
RA       You M., Ku P.-T., Hedlickova R., Bone H.R., Jr.;
RF       MEDLINE=960360; PubMed=937286; bone H.R., Jr.;
AT       mediator, a member of the inhibitor-of-apoptosis protein family, is a
AT       member of the antiapoptotic activity of the v-rel oncoprotein." ;
RL       Mol. Cell. Biol. 17:7328-7341(1997).
CC       -1 FUNCTION: SUPPRESSOR OF APOPTOSIS IN ONCOPROTEIN V-REL-TRANSFORMED
CC       CELLS.
CC       -1 SUBCELLULAR LOCATION: EXPRESSED PREDOMINANTLY IN THE CYTOPLASM OF
CC       THE V-REL-TRANSFORMED CELLS.
CC       -1 THE SPECIFIC mRNA IS EXPRESSED AT RELATIVELY HIGH LEVELS IN THE
CC       SPINAL MUSCLE AND SKELTERAL LONG, AND AT VERY LOW LEVELS
CC       IN TESTIS, BRAIN AND SKLETERAL MUSCLE.
CC       -1 INDUCTION: INDUCED DURING THE V-REL-MEDIATED TRANSFORMATION
CC       PROCESS.
CC       -1 DOMAIN: THE RING FINGER IS IMPORTANT FOR ITS ANTIAPOPTOTIC EFFECT.
CC       -1 SIMILARITY: MEMBER OF THE IAP FAMILY.
CC       -1 SIMILARITY: CONTAINS 3 BIR DOMAINS (BACULOVIRAL INHIBITION OF

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CC APOPTOSIS PROTEIN REPEAT.)
CC -1 SIMILARITY: CONTAINS A C3HC4-CLASS ZINC FINGER.
DR EMBL: AF0089592; AAB88044.1; -.
DR HSPB: Q13490; IGBH.
DR InterPro: IPRO01370; BIR.
DR InterPro: IPRO01375; CARD.
DR InterPro: IPRO01411; CAT_Lfing.
DR Pfam: PF00653; BIR_3.
DR Pfam: PF00097; zfcCHC4; 1.
DR SMART: SMO0238; BIR; 3.
DR SMART: SMO0114; CARD; 1.
DR PROSITE: PS01282; BIR_REPEAT_1; 3.
DR PROSITE: PS0143; BIR_REPEAT_2; 3.
DR PROSITE: PS0209; CARD; 1.
DR Consists: L3c-Lfing; Repeat.
DR REPEAT 1.
FT REPEAT 176 242 BIR_REPEAT_1.
FT REPEAT 262 329 BIR_REPEAT_2.
FT ZN_FING 563 597 C3HC4-TYPE.
SO SEQUENCE 610 AA; 68924 MW; ADP4761965084A6 CRC64; 50

Query Match          98.3%; Score 290; DB 13; Length 610;
Best Local Similarity 95.8%; Pred. No. 2e-29; 1; Indexes 0; Gaps 0;
Matches 46; Conservative 1; Mismatches 1;

Cy      1  PRLDASGFFYGRNDVCKPCGGSLKRCNSDDDPWEAKKPFRC 48
Db      280 PRLADAGFYFGRNDVKCFCCCGSLRNCSSSDPWIEHAKWPFRC 327
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
RESULT    4
Q90NH1     OG0NH1     PRELIMINARY:   PRt: 1140 AA.
ID         OG0NH1     :
RA         OG0NH1     :
DT         01-MAY-2000 (TREMblrel. 13, Created)
DI         01-MAY-2000 (TREMblrel. 13, Last sequence update)
DT         01-JUN-2001 (TREMblrel. 17, Last annotation update)
DE         AP12-MLT Fusion PROTEIN.
GN         AP12-MLT.
GC         Homo sapiens (Human).
OC         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC         Mamalia; Eutheria; Primates; Catarrhini; Hominiidae; Homo.
OX         NCSTX_textid=9606;
RX         MEDLINE=93272420; PubMed=10339644;
RA         Dietlamm J., Beems M., Wlodarski I., Stefanova-Ouzounova M.,
RA         Hernandez J.N., Hosfeld D.K., De Wolf-Petersen C., Hagemeier A.,
RA         Van den Berghe H., Maynen P.:
RA         "The apoptosis inhibitor gene AP12 and a novel 18q gene, MLT, are
RT         recurrently rearranged in the t(11;18)(q21;q21)p65associated with
RT         mucosa-associated lymphoid tissue lymphomas."
RT         Blood 93:3601-3609(1999).
KC         -1 SIMILARITY TO IMMUNOGLOBULIN AND MAJOR HISTOCOMPATIBILITY COMPLEX
CC EMBL: AF123054; MAD6161.1; -.
DR HSPB: Q13490; IOBH.
DR InterPro: IPRO01370; BIR.
DR InterPro: IPRO01375; Caspase.
DR InterPro: IPRO01309; ICE_p20.
DR InterPro: IPRO03598; 19_c2.
DR InterPro: IPRO03600; 19_1Mc.
DR InterPro: IPRO03006; 19_Mhc.
DR Pfam: PF00653; BIR; 3.
DR Pfam: PF00653; BIR_3.
DR SMART: SMO0238; BIR; 3.
DR SMART: SMO0115; CASC; 1.
DR SMART: SMO0408; IGc2; 1.
DR SMART: SMO0410; IGc-like; 1.
DR PROSITE: PS01282; BIR_REPEAT_1; 3.
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DR PROSITE: PS50143; BIR\_REPEAT\_2; 3.  
DR PROSITE: PS50209; CASPASE\_P20; 1.  
SQ SEQUENCE 1140 AA; 128738 MW; 0C18D90287C723E CRC64;

Query Match 95.6%; Score 282; DB 4; Length 1140;  
Best Local Similarity 93.8%; Pred. No. 4.2e-28;  
Matches 45; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

OY 1 PEOLASAGFYVGRNDVYKCFCCDGLRCWESGDDPVEHAKWFPCE 48  
DB 273 PEOLASAGFYVGRNDVYKCFCCDGLRCWESGDDPVEHAKWFPCE 320

RESULT 5  
OQ9ZC6 PRELIMINARY; PRT; 589 AA.

DT 01-MAY-2000 (TREMBLrel. 13, Created)  
DT 01-MAY-2000 (TREMBLrel. 13, Last sequence update)  
DT 01-JUN-2001 (TREMBLrel. 17, Last annotation update)  
DE INHIBITOR OF APOPTOSIS PROTEIN 2.  
OS Rattus norvegicus (Rat).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.  
OX NCBI\_TaxID=10116;  
RN [1]

RP SEQUENCE FROM N.A.  
RC TISSUE=KIDNEY;  
RA Dong Z., Denton M., Gu S.M., Saikumar P., Venkatachalam M.A.;  
RT "Cloning of cDNA for rat inhibitor of apoptosis protein 2.";  
RL Submitted (SEP-1999) to the EMBL/Genbank/DBJ databases.  
-1- SIMILARITY: CONTAINS A RING-TYPE ZINC FINGER.

DR EMBL: AF190020; AAF04585.1;  
DR HSSP: Q13490; 108B;  
DR InterPro: IPR001370; BIR.  
DR InterPro: IPR001315; CARD.  
DR InterPro: IPR001841; ZnF\_ring.  
DR Pfam: PF00653; BIR; 3.  
DR Pfam: PF00619; CARD; 1.  
DR Pfam: PF00097; ZF-C3HC4; 1.  
DR SMART: SM00238; BIR; 3.  
DR SMART: SM00114; CARD; 1.  
DR SMART: SM00184; RING; 1.  
DR PROSITE: PS01282; BIR\_REPEAT\_1; 3.  
DR PROSITE: PS50143; BIR\_REPEAT\_2; 3.  
DR PROSITE: PS50209; CARD; 1.  
KW Zinc-finger  
SQ SEQUENCE 589 AA; 66777 MW; E6812FE3EA34142 CRC64;

Query Match 93.9%; Score 277; DB 11; Length 589;  
Best Local Similarity 91.7%; Pred. No. 9.4e-28;  
Matches 44; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

OY 1 PEOLASAGFYVGRNDVYKCFCCDGLRCWESGDDPVEHAKWFPCE 48  
DB 259 PEOLASAGFYVGRNDVYKCFCCDGLRCWESGDDPVEHAKWFPCE 306

RESULT 6  
OQ9SE8 PRELIMINARY; PRT; 589 AA.

DT 01-MAR-2001 (TREMBLrel. 16, Created)  
DT 01-MAR-2001 (TREMBLrel. 16, Last sequence update)  
DT 01-JUN-2001 (TREMBLrel. 17, Last annotation update)  
DE INHIBITOR OF APOPTOSIS PROTEIN 2.  
OS Rattus norvegicus (Rat).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.  
OX NCBI\_TaxID=10116;  
RN [1]

RP SEQUENCE FROM N.A.  
RA Holcik M., Lefebvre C.A., Hicks K., Korneluk R.G.;  
RT "Cloning and Characterization of the Rat Homologs of the Inhibitor of  
RT Apoptosis Protein 1, 2, and 3 Genes."  
RL Submitted (SEP-1999) to the EMBL/Genbank/DBJ databases.  
CC -1- SIMILARITY: CONTAINS A RING-TYPE ZINC FINGER.

DR EMBL: AF18431; AAG2971.1;  
DR InterPro: IPR001370; BIR.  
DR InterPro: IPR001315; CARD.  
DR InterPro: IPR001841; ZnF\_ring.  
DR Pfam: PF00653; BIR; 3.  
DR Pfam: PF00619; CARD; 1.  
DR Pfam: PF00097; ZF-C3HC4; 1.  
DR SMART: SM00238; BIR; 3.  
DR SMART: SM00114; CARD; 1.  
DR SMART: SM00184; RING; 1.  
DR PROSITE: PS01282; BIR\_REPEAT\_1; 1.  
DR PROSITE: PS50143; BIR\_REPEAT\_2; 3.  
DR PROSITE: PS50209; CARD; 1.  
KW Zinc-finger.  
SQ SEQUENCE 589 AA; 66750 MW; BAF7089BD7CD285B CRC64;

Query Match 93.9%; Score 277; DB 11; Length 589;  
Best Local Similarity 91.7%; Pred. No. 9.4e-28;  
Matches 44; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

OY 1 PEOLASAGFYVGRNDVYKCFCCDGLRCWESGDDPVEHAKWFPCE 48  
DB 259 PEOLASAGFYVGRNDVYKCFCCDGLRCWESGDDPVEHAKWFPCE 306

RESULT 7  
OQ9SE9 PRELIMINARY; PRT; 602 AA.

DT 01-MAR-2001 (TREMBLrel. 16, Created)  
DT 01-MAR-2001 (TREMBLrel. 16, Last sequence update)  
DT 01-JUN-2001 (TREMBLrel. 17, Last annotation update)  
DE INHIBITOR OF APOPTOSIS PROTEIN 1.  
OS Rattus norvegicus (Rat).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.  
OX NCBI\_TaxID=10116;  
RN [1]

RP SEQUENCE FROM N.A.  
RA Holcik M., Lefebvre C.A., Hicks K., Korneluk R.G.;  
RT "Cloning and Characterization of the Rat Homologs of the Inhibitor of  
RT Apoptosis Protein 1, 2, and 3 Genes."  
RL Submitted (SEP-1999) to the EMBL/Genbank/DBJ databases.  
CC -1- SIMILARITY: CONTAINS A RING-TYPE ZINC FINGER.

DR EMBL: AF18430; AAG2970.1;  
DR InterPro: IPR001370; BIR.  
DR InterPro: IPR001315; CARD.  
DR InterPro: IPR001841; ZnF\_ring.  
DR Pfam: PF00653; BIR; 3.  
DR Pfam: PF00619; CARD; 1.  
DR Pfam: PF00097; ZF-C3HC4; 1.  
DR SMART: SM00238; BIR; 3.  
DR SMART: SM00114; CARD; 1.  
DR SMART: SM00184; RING; 1.  
DR PROSITE: PS01282; BIR\_REPEAT\_1; 1.  
DR PROSITE: PS50143; BIR\_REPEAT\_2; 3.  
DR PROSITE: PS50209; CARD; 1.  
KW Zinc-finger.  
SQ SEQUENCE 602 AA; 67326 MW; CC91385FEA62DDE5A CRC64;

Query Match 93.2%; Score 275; DB 11; Length 602;  
Best Local Similarity 89.6%; Pred. No. 1.7e-27;  
Matches 43; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

OY 1 PEOLASAGFYVGRNDVYKCFCCDGLRCWESGDDPVEHAKWFPCE 48

Db 273 PDLASAGFYVGHSDVYKCFCCDGLRCMESGDDPWVEMHAKMPFCE 320

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RESULT 8
ID 09DND2 PRELIMINARY; PRT: 324 AA.
AC 09DND2;
DT 01-MAR-2001 (TREMBlrel. 16, Created)
DT 01-MAR-2001 (TREMBlrel. 16, Last sequence update)
DT 01-JUN-2001 (TREMBlrel. 17, Last annotation update)
DE APOPTOSIS INHIBITOR CH-1AP1 (FRAGMENT).
OS Gallus gallus (Chicken)
OC Archaeoptera; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
NC Eukaryota; Metazoa; Neognathae; Galliformes; Phasianidae; Phasianinae;
GX NCBI_TaxID=9031;
RN 111
RS SEQUENCE FROM N.A.
RA Penhelen C.N., Bargmann W.J., Varadarajan J., Bose H.R. Jr.;
RT The apoptosis inhibitor ch-1ap1 is a direct transcriptional target of
RL EMBL AF011280: AAC42316.1; -.
DR EMBL AF011280: AAC42316.1; -.
DR InterPro: IPR001370: BIR.
DR Pfam: PF00653: BIR. 3.
DR SMART: SM00238: BIR. 3.
DR PROSITE: PS01282: BIR_REPEAT_1: 1.
DR PROSITE: PS01443: BIR_REPEAT_2: 3.
FT NON_TER 324
SC SEQUENCE 324 AA: 36567 MW: 58289DEAE373P3 CRC64:

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Query Match 91.9%; Score 271; DB 13; Length 324;  
 Best Local Similarity 95.6%; Pred. No. 36-27; Indels 0; Gaps 0;  
 Matches 43; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

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Qy 1 PDLASAGFYVGRNDVYKCFCCDGLRCMESGDDPWVEMHAKMPF 45
Db 280 PDLASAGFYVGRNDVYKCFCCDGLRCMESGDDPWVEMHAKMPF 324

RESULT 9
ID 088642 PRELIMINARY; PRT: 224 AA.
AC 088642;
DT 01-NOV-1998 (TREMBlrel. 08, Created)
DT 01-NOV-1998 (TREMBlrel. 08, Last sequence update)
DT 01-JUN-2001 (TREMBlrel. 17, Last annotation update)
DE INHIBITOR OF APOPTOSIS PROTEIN (FRAGMENT).
CN RIAP1.
OS Rattus norvegicus (Rat).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
NC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
GX NCBI_TaxID=10116;
RN 111
RS SEQUENCE FROM N.A.
RA STRAIN-WISTAR; TISSUE-Ovary, CORPUS LUTEUM;
RA Bradley C.K., Lareu R.R., Dharmarajan A.M.;
RT Cloning and characterization of an inhibitor of apoptosis protein
RT (IAP) in the rat corpus luteum.
RL Submitted (Aug-1998) to the EMBL/GenBank/DBJ databases.
DR EMBL AF081503: AAC32497.1; -.
DR HSP: Q13490: IGBR.
DR InterPro: IPR001370: BIR.
DR Pfam: PF00653: BIR. 2.
DR SMART: SM00238: BIR. 2.
DR PROSITE: PS01443: BIR_REPEAT_2: 3.
FT NON_TER 224
SC SEQUENCE 224 AA: 25209 MW: 213A52534D5E56A CRC64:

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Query Match 70.8%; Score 209; DB 11; Length 224;  
 Best Local Similarity 91.9%; Pred. No. 2,26-19;  
 Matches 34; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 1 PDLASAGFYVGRNDVYKCFCCDGLRCMESGDDPWVEMHAKMPF 45  
 Db 188 PDLASAGFYVGRNDVYKCFCCDGLRCMESGDDPWVEMHAKMPF 224

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RESULT 10
ID 09EN27 PRELIMINARY; PRT: 264 AA.
AC 09EN27;
DT 01-MAR-2001 (TREMBlrel. 16, Created)
DT 01-MAR-2001 (TREMBlrel. 16, Last sequence update)
DT 01-JUN-2001 (TREMBlrel. 17, Last annotation update)
DE AAV021.
GN AAV021.
OS Anaseta moorei entomopoxvirus (AmePV).
OC Viruses; dsDNA viruses, no RNA stage; Poxvirinae; Entomopoxvirinae;
NC Entomopoxvirus B.
GX NCBI_TaxID=28321;
RN 111
RS SEQUENCE FROM N.A.
RA MEDLINE=20396580; PubMed=10936094;
RA Bowden A.L., Glassberg K.J., Digans J., Shaw R., Farmerie W.,
RT Complete genomic sequence of the Anaseta moorei Entomopoxvirus:
RT Analysis and comparison with other Poxviruses.
RT Virology 274:120-139(2000).
FT [2]
RN 121
RS SEQUENCE FROM N.A.
RA Bowden A.L., Glassberg K.J., Digans J., Shaw R., Farmerie W.,
RA Meyer R.W.;
RL Submitted (MAR-2000) to the EMBL/GenBank/DBJ databases.
DR EMBL AF20284: AAC02727.1; -.
DR InterPro: IPR001370: BIR.
DR InterPro: IPR001841: ZnF_Ring.
DR Pfam: PF00653: BIR. 2.
DR SMART: SM00238: BIR. 2.
DR SMART: SM00184: RING. 1.
DR PROSITE: PS01443: BIR_REPEAT_2: 2.
SC SEQUENCE 264 AA: 30547 MW: 28B72DA4B58D920A CRC64:

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Query Match 67.1%; Score 198; DB 12; Length 264;  
 Best Local Similarity 61.7%; Pred. No. 76-18;  
 Matches 29; Conservative 10; Mismatches 8; Indels 0; Gaps 0;

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Qy 2 PDLASAGFYVGRNDVYKCFCCDGLRCMESGDDPWVEMHAKMPFCE 48
Db 126 ERLASAGFYVGRNDVYKCFCCDGLRCMESGDDPWVEMHAKMPFCE 172

RESULT 11
ID 09HAP7 PRELIMINARY; PRT: 280 AA.
AC 09HAP7;
DT 01-MAR-2001 (TREMBlrel. 16, Created)
DT 01-MAR-2001 (TREMBlrel. 16, Last sequence update)
DT 01-JUN-2001 (TREMBlrel. 17, Last annotation update)
DE LIVIN INHIBITOR OF APOPTOSIS (INHIBITOR OF APOPTOSIS) (HA261N11.1.1)
DE (BACULOVIRAL IAP REPEAT-CONTAINING PROTEIN 7 (LIVIN), ISOFORM 1).
GN LIVIN OR BIRC7.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
NC Mammalia; Eutheria; Primates; Catartini; Hominoidea; Homo.
GX NCBI_TaxID=9606;
RN 111
RS SEQUENCE FROM N.A.
RA Kasof G.M., Gomes B.C.;
RT "Livin, a novel inhibitor of apoptosis (IAP) family member.":
RL J. Biol. Chem. 0:0-0(2000).

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RN [2]
RP SEQUENCE FROM N.A.
RA Ashbald I., Allan A., Pollack A., Panet A., Ben-Yehuda D.;
RT different biological properties and tissue distribution pattern.*;
RL Submitted (Apr-2001) to the EMBL/GenBank/DBJ databases.
RN [3]
RP SEQUENCE FROM N.A.
RA Hall R.;
RL Submitted (Apr-2001) to the EMBL/GenBank/DBJ databases.
CC -1- SIMILARITY: CONTAINS A RING-TYPE ZINC FINGER.
DR EMBL: AF311388; AAC35522.1; -;
DR EMBL: AJ309298; CAC37337.1; -;
DR EMBL: AL216277; CAC36112.1; -;
DR InterPro: IPR001370; BIR;
DR InterPro: IPR001841; ZnF_ring.
DR Pfam: PF00653; BIR; 1.
DR SMART: SM00387; ZF-C3HC4; 1.
DR SMART: SM00184; RING; 1.
DR PROSITE: PS50143; BIR_REPEAT_2; 1.
DR PROSITE: PS00518; ZINC_FINGER_C3HC4; UNKNOWN_1.
DR Zinc-finger.
KW SEQUENCE 280 AA; 30866 MW; 630BE9C0737F7952 CRC64;
SQ

Query Match 66.1%; Score 195; DB 4; Length 280;
Best Local Similarity 62.5%; Pred. No. 1,9e-17;
Matches 30; Conservative 7; Mismatches 11; Indels 0; Gaps 0;

OY 1 PEOLASAGFTYVGNDYKCCGCGGLRCWESQDDPVYHAKWPFRC 48
DB 105 PELLAAGFFRHGDYKCFYGGGLQSWKRRDDPWTEHAKWPFSCQ 152

RESULT 12
ID Q9H2A8 PRELIMINARY; PRT; 298 AA.
AC Q9H2A8;
DT 01-MAY-2001 (TrEMBLrel. 16, created)
DT 01-MAY-2001 (TrEMBLrel. 16, last sequence update)
DT 01-JUN-2001 (TrEMBLrel. 17, last annotation update)
DE INHIBITOR OF APOPTOSIS PROTEIN KIN (BAX61M1.1.2) (BACULOVIRAL IAP
DE REPEAT-CONTAINING PROTEIN 7 (LITVIN), ISOFORM 2).
GN BIRC7.
OS Homo sapiens (human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Homiidae; Homo.
OX NCBI_Taxid=9606;
RN [1]
RP SEQUENCE FROM N.A.
RA Tissue-Kinney;
RC Lin J.-H., Peng G., Huang Q., Morser J.;
RT A novel member of the inhibitor of apoptosis protein family.*;
RL Biochem. Biophys. Res. Commun. 0:0-0(2000).
RN [2]
RP SEQUENCE FROM N.A.
RA Ashbald I., Allan A., Pollack A., Panet A., Ben-Yehuda D.;
RT different biological properties and tissue distribution pattern.*;
RL Submitted (Apr-2001) to the EMBL/GenBank/DBJ databases.
RN [3]
RP SEQUENCE FROM N.A.
RA Hall R.;
RL Submitted (Apr-2001) to the EMBL/GenBank/DBJ databases.
CC -1- SIMILARITY: CONTAINS A RING-TYPE ZINC FINGER.
DR EMBL: AF301009; AAC37878.1; -;
DR EMBL: AJ309298; CAC37338.1; -;
DR EMBL: AL216277; CAC36111.1; -;
DR InterPro: IPR001370; BIR;
DR InterPro: IPR001841; ZnF_ring.
DR Pfam: PF00653; BIR; 1.
DR Pfam: PF00097; ZF-C3HC4; 1.
SQ

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DR SMART: SM00238; BIR; 1.
DR SMART: SM00184; RING; 1.
DR PROSITE: PS50143; BIR_REPEAT_2; 1.
DR PROSITE: PS00518; ZINC_FINGER_C3HC4; UNKNOWN_1.
DR Zinc-finger.
KW SEQUENCE 298 AA; 32798 MW; B2BAEE531BECD1 CRC64;
SQ

Query Match 66.1%; Score 195; DB 4; Length 298;
Best Local Similarity 62.5%; Pred. No. 1,9e-17;
Matches 30; Conservative 7; Mismatches 11; Indels 0; Gaps 0;

OY 1 PEOLASAGFTYVGNDYKCCGCGGLRCWESQDDPVYHAKWPFRC 48
DB 105 PELLAAGFFRHGDYKCFYGGGLQSWKRRDDPWTEHAKWPFSCQ 152

RESULT 13
ID Q9YNL8 PRELIMINARY; PRT; 281 AA.
AC Q9YNL8;
DT 01-MAY-1999 (TrEMBLrel. 10, created)
DT 01-MAY-1999 (TrEMBLrel. 10, last sequence update)
DT 01-JUN-2001 (TrEMBLrel. 17, last annotation update)
DE IAP PROTEIN.
GN IAP.
OS Choriostoneura fumiferana nuclear polyhedrosis virus (CFMNPV).
OC Viruses; dsDNA viruses, no RNA stage; Baculoviridae;
OC Nucleopolyhedrovirus.
OX NCBI_Taxid=10448;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=IRELAND;
RA Lawton H., Ait B.M., Ladd T., Pall R.;
RC Cloned (Genbank) to the EMBL/GenBank/DBJ databases.
CC -1- SIMILARITY: CONTAINS A RING-TYPE ZINC FINGER.
DR EMBL: L482510; IAP00537.1; -;
DR HSP: 013490; IAP;
DR InterPro: IPR001370; BIR;
DR InterPro: IPR001841; ZnF_ring.
DR Pfam: PF00653; BIR; 2.
DR Pfam: PF00097; ZF-C3HC4; 1.
DR SMART: SM00238; BIR; 2.
DR SMART: SM00184; RING; 1.
DR PROSITE: PS01262; BIR_REPEAT_1; UNKNOWN_1.
DR PROSITE: PS50143; BIR_REPEAT_2; 2.
DR Zinc-finger.
KW SEQUENCE 281 AA; 32090 MW; B2D9BE8A359F105F CRC64;
SQ

Query Match 65.1%; Score 192; DB 12; Length 281;
Best Local Similarity 60.0%; Pred. No. 4,5e-17;
Matches 31; Conservative 5; Mismatches 11; Indels 0; Gaps 0;

OY 1 PEOLASAGFTYVGNDYKCCGCGGLRCWESQDDPVYHAKWPFRC 47
DB 142 PELLAAGFFRHGDYKCFYGGGLQSWKRRDDPWTEHAKWPFSCQ 188

RESULT 14
ID G9U492 PRELIMINARY; PRT; 379 AA.
AC G9U492;
DT 01-MAY-2000 (TrEMBLrel. 13, created)
DT 01-MAY-2000 (TrEMBLrel. 13, last sequence update)
DT 01-JUN-2001 (TrEMBLrel. 17, last annotation update)
DE INHIBITOR OF APOPTOSIS PROTEIN.
GN IAP1.
OS Trichoplusia ni (cabbage looper).
OC Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
OC Pterygota; Neoptera; Endopterygota; Lepidoptera; Glossata; Ditylata;
OC Noctuidae; Noctuidae; Plusiinae; Trichoplusia.
SQ

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Search completed: January 7, 2002, 16:03:54  
Job time: 1413 sec

OK NCBI\_TaxID=7111;  
RN 11)  
RP SEQUENCE FROM N.A.  
RA MEDLINE=20082908; PubMed=10593985;  
RA Sesnagiri S., Vucic D., Lee J., Dixit V.M.;  
RT Baculovirus-based genetic screen for antiapoptotic genes identifies a  
RT novel IAP.  
RL J. Biol. Chem. 274:36769-36773(1999).  
DR EMBL: AF195528; AAF19619.1; .  
DR HSSP: Q13490; 109H.  
DR InterPro: IPR01370; BIR.  
DR InterPro: IPR01841; ZnF\_Ling.  
DR Pfam: PF00523; BIR\_2.  
DR SMART: SM00238; BIR\_2.  
DR PROSITE: PS01282; BIR\_REPEAT\_1; UNKNOWN\_2.  
DR PROSITE: PS0143; BIR\_REPEAT\_2.  
SO SEQUENCE 379 AA; 41857 MW; 069381A012D9D65 CRC64;

Query Match 64.1%; Score 189; DB 5; Length 379;  
Best Local Similarity 61.7%; Pred. No. 1,5e-16;  
Matches 29; Conservative 7; Mismatches 11; Indels 0; Gaps 0;

OY 1 PROLASGFFYVGRNDVRCGCCGILRCMSGDDPWEHAKWPPRC 47  
DB 227 PEEIAXGFFYTGCGDKTKCFYCDSGLKMDNDVPEWQHAWPDR 273

## RESULT 15

OGNJ07 PRELIMINARY: PRT; 377 AA.  
AC OGNJ07.  
DT 01-OCT-2000 (TRENBERG, 15, Created)  
DT 01-OCT-2000 (TRENBERG, 15, Last sequence update)  
DE 01-JUN-2001 (TRENBERG, 17, Last annotation update)  
DE INHIBITOR OF APOPTOSIS PROTEIN.  
GN IAP.  
OS Spodoptera frugiperda (Fall armyworm).  
OC Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;  
OC Pterygota; Neoptera; Endopterygota; Lepidoptera; Glossata; Ditylsia;  
OC Noctuidae; Noctuidae; Amphipyritae; Spodoptera.  
OX NCBI\_TaxID=7108;  
RN 11)  
RP SEQUENCE FROM N.A.  
RA MEDLINE=2014055; PubMed=10677478;  
RA Huang O., Devereux O.L., Maeda S., Salvanes G.S., Stennicke H.R.,  
RA Hammock B.D., Reed J.C.;  
RT "Evolutionary conservation of apoptosis mechanisms: Lipidopteran and  
RT baculoviral inhibitor of apoptosis proteins are inhibitors of  
RT mammalian caspase-9."  
RL Proc. Natl. Acad. Sci. U.S.A. 97:1427-1432(2000).  
DR EMBL: AF186378; AAF5285.1; .  
DR InterPro: IPR01370; BIR.  
DR InterPro: IPR01841; ZnF\_Ling.  
DR Pfam: PF00523; BIR\_2.  
DR SMART: SM00238; BIR\_2.  
DR PROSITE: PS01282; BIR\_REPEAT\_1; UNKNOWN\_2.  
DR PROSITE: PS0143; BIR\_REPEAT\_2.  
SO SEQUENCE 377 AA; 41871 MW; F1207D2D3ED0B38 CRC64;

Query Match 63.7%; Score 188; DB 5; Length 377;  
Best Local Similarity 61.7%; Pred. No. 2e-16;  
Matches 29; Conservative 7; Mismatches 11; Indels 0; Gaps 0;

OY 1 PROLASGFFYVGRNDVRCGCCGILRCMSGDDPWEHAKWPPRC 47  
DB 228 PEEIAXGFFYTGCGDKTKCFYCDSGLKMDNDVPEWQHAWPDR 274

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